

WEST

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L16: Entry 1 of 19

File: USPT

DOCUMENT-IDENTIFIER: US 6403315 B1

TITLE: Method and apparatus for DNA sequencing and DNA identification

Detailed Description Text (123):

Using the described theoretical principles as a guide for experiments, reliable hybridizations have been obtained with probes six to eight nucleotides in length. All experiments were performed with a floating plastic sheet providing a film of hybridization solution above the filter. This procedure allows maximal reduction in the amount of probe, and thus reduced label costs in dot blot hybridizations. The high concentration of sodium lauroyl sarcosine instead of sodium lauroyl sulfate in the phosphate hybridization buffer allows dropping the reaction from room temperature down to 12.degree. C. Similarly, the 4-6.times.SSC, 10% sodium lauroyl sarcosine buffer allows hybridization at temperatures as low as 2.degree. C. The detergent in these buffers is for obtaining tolerable background with up to 40 nM concentrations of labelled probe. Preliminary characterization of the thermal stability of short oligonucleotide hybrids was determined on a prototype octamer with 50% G+C content, i.e. probe of sequence TGCTCATG. The theoretical expectation is that this probe is among the less stable octamers. Its transition enthalpy is similar to those of more stable heptamers or, even to probes 6 nucleotides in length (Bresslauer et al., Proc. Natl. Acad. Sci. U.S.A. 83: 3746 (1986)). Parameter T.sub.d, the temperature at which 50% of the hybrid is melted in unit time of a minute is 18.degree. C. The result shows that T.sub.d is 15.degree. C. lower for the 8 bp hybrid than for an 11 bp duplex (Wallace et al., Nucleic Acids Res. 6: 3543 (1979)).

W0 00

The Method

M. L. Jones

1/15/92

Set Name Query

side by side

Hit Count Set Name

result set

DB=USPT,JPAB,EPAB,DWPI; PLUR=YES; OP=ADJ

<u>L17</u>	breslauer	228	<u>L17</u>
<u>L16</u>	breuslauer	0	<u>L16</u>
<u>L15</u>	enthalpy same nucleic same acid same duplex	19	<u>L15</u>
<u>L14</u>	screen\$ adj thermostab\$ same equil\$ same duplex	0	<u>L14</u>
<u>L13</u>	screen\$ same thermostab\$ same equil\$ same duplex	0	<u>L13</u>
<u>L12</u>	thermostab\$ same (mismatch\$ or exocyclic or polymorph\$) same duplex same thermal	0	<u>L12</u>
<u>L11</u>	screen\$ same nucleic\$ same acid same duplex\$ same stability adj label\$ adj energ\$	0	<u>L11</u>
<u>L10</u>	screen\$ same nucleic\$ same acid same duplex\$ same stability adj label\$ adj FET	0	<u>L10</u>
<u>L9</u>	screen\$ same nucleic\$ same acid\$ same duplex\$ same stability adj label\$ adj FET	0	<u>L9</u>
<u>L8</u>	screen\$ same nucleic same acid same duplex same stability adj label\$ adj FET	0	<u>L8</u>
<u>L7</u>	screen\$ same nucleic same acid same duplex same stability same label\$ same FET	1	<u>L7</u>
<u>L6</u>	screen\$ same nucleic same acid same duplex same stability same label\$	7	<u>L6</u>
<u>L5</u>	screen\$ same nucleic same acid same duplex same stability and equil\$	7	<u>L5</u>
<u>L4</u>	screen\$ same nucleic same acid same duplex same stability and equilibra	0	<u>L4</u>
<u>L3</u>	screen\$ same nucleic same acid same duplex same stability adj equilibra	0	<u>L3</u>
<u>L2</u>	screen\$ same nucleic same acid same duplex same stability same equilibra	0	<u>L2</u>
<u>L1</u>	screen\$ same nucleic same acid same duplex same stability	13	<u>L1</u>

END OF SEARCH HISTORY

Maupin, Christin

To: STIC-ILL
Subject: screening nuclei acid duplex stability 09/869004
Sensitivity: Private

- Lisitsyn et al, Cloning the differences between two complex genomes, *Science*, 259: 946-951 (1993).
- Koss et al, "Flow cytometric measurements of DNA and other cell components in human tumors: a critical appraisal," *Human Pathology*, 20: 528-548 (1989).
- Van Ness et al, "A versatile solid support system for oligodeoxynucleotide probe-based hybridization assays," *Nucleic Acids Research*, 19: 3345-3350 (1991).
- Fulton et al, "Advanced multiplexed analysis iwth the FlowMetrix.TM. system," *Clinical Chemistry*, 43: 1749-1756 (1997).
- Hakala et al, "Time-resolved fluorescence detection of oligonucleotide hybridization on a single microparticle: covalent immobilization of oligonucleotides and quantitation of a model system," *Bioconjugate Chemistry*, 8: 232-237 (1997).
- Hakala et al, "Detection of oligonucleotide hybridization on a single microparticle by time-resolved fluorometryL hybridization assays on polymer particles obtained by direct solid phase assembly of the oligonucleotide probes," *Bioconjugate Chemistry*, 8: 378-384 (1997).
- Raineri et al, "Improved efficiency for single-sided PCR by creating a reusable pool of first-strand cDNA coupled to a solid phase," *Nucleic Acids Research*, 19:4010 (1991).
- Lee et al, "Reusable cDNA libraries coupled to magnetic beads," *Anal. Biochem.*, 206: 206-207 (1992).
- Lovgren et al, "Sensitive bioaffinity assays with individual microparticles and time-resolved fluorometry," *Clinical Chemistry*, 43: 1937-1943 (1997).
- Product brochure, "Dynabeads template preparation kit," Dynal Inc. (Great Neck, New York), 1991.
- Vlieger et al, "Quantitation of polymerase chain reaction products by hybridizaton-based assays with fluorescent, colorimetric, or chemiluminscent detection," *Anal. Biochem.*, 205: 1-7 (1992).
- Khudyakov et al, "Primer specific solid-phase detection of PCR products," *Nucleic Acids Research*, 22: 1320-1321 (1994).
- Wan et al, "Cloning differentially expressed mRNAs," *Nature Biotechnology*, 14: 1685-1691.

Stamm et al, "Sanchoed PCR: PCR with cDNA coupled to a solid phase," *Nucleic Acids Research*, 19: 1350 (1991).

Beattie et al, "Advances in genosensor research," *Clin. Chem.*, 41(5): 700-706 (1995).

Sagerstrom et al, "Subtractive cloning: past, present, and future," *Annu. Rev. Biochem.*, 66: 751-783 (1997).

Yershov et al, "DNA analysis and diagnostics on oligonucleotide microchips," *Proc. Natl. Acad. Sci.*, 93: 4913-4918 (1996).

Chee et al, "Accessing genetic information with high-density DNA arrays," *Science*, 274: 610-614 (1996).

Lipshutz et al, "Using oligonucleotide probe arrays to access genetic diversity," *BioTechniques*, 19: 442-447 (1995).

Shalon et al, "A DNA microarray system for analyzing complex DNA samples using two-color fluorescent probe hybridization," *Genome Research*, 6: 639-645 (1996).

Schena et al, "Quantitative monitoring of gene expression patterns with a complementary DNA microarray," *Science*, 270:467-470 (1995).

DeRisi et al, "Exploring the metabolic and genetic control of gene expression on a genomic scale," *Science*, 278: 680-686 (1997).

Zhao et al, "High-density cDNA filter analysis: a novel approach for large-scale, quantitative analysis of gene expression," *Gene*, 156: 207-213 (1995).

Wodicka et al, "Genome-wide expression monitoring in *Saccharomyces cerevisiae*," *Nature Biotechnology*, 15: 1359-1367(1997).

Shoemaker et al, "Quantitative phenotypic analysis of yeast deletion mutants using a highly parallel molecular bar-coding strategy," *Nature Genetics*, 14: 450-456 (1996).

Unrau et al, "Non-cloning amplification of specific DNA fragments from whole genomic DNA digests using DNA indexers," *Gene*, 145: 163-169 (1994).

Broude et al, "Enhanced DNA sequencing by hybridization," *Proc. Natl. Acad. Sci.*, 91: 3072-3076 (1994).

Hultman et al, "Direct solid phase sequencing of genomic and plasmid DNA using magnetic beads as solid support," *Nucleic Acids Research*, 17: 4937-4946 (1989).

Hensel et al, "Simultaneous identification of bacterial virulence genes by negative selection," *Science*, 269: 400-403 (1995).

Needels et al, "Generation and screening of an oligonucleotide-encoded synthetic peptide library," *Proc. Natl. Acad. Sci.*, 90: 10700-10704 (1993).

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